FIG. 1B

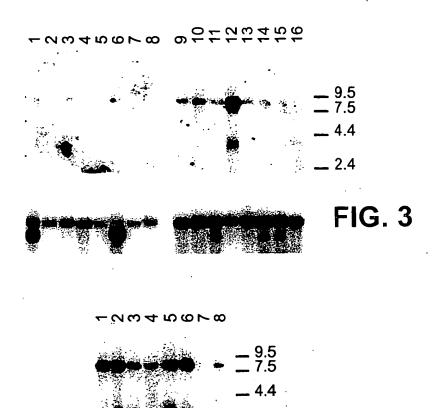
RKKGDAEN---MVLCDGCDRGHHTYCV--RPKLKTVPEGD-WFCPEC --MIVCDGCESGCHMECF--RPRMTKVPEGD-WFCQRC --RGNNEDK---LLLCDGCDDSYHTFCL--IPPLPDVPKGD-WRCPKC -TKEQNREKKPEELISCADCGNSGHPSCLKFSPELTVRVKALRWQCIEC ----QGKNADN---MLFCDSCDRGFHMECC--DPPLTRMPKGM-WICQIC CEKCFNEIQGESVSLGDDPSQPQTTINKEQFSKRKNDTLDPELFVECTECGRKMHQICV--LHHEIIWPAG--FVCDGC CEKCFTEIQGENVTLGDDPSQPQTTISKDQFE**KKKND**TLD**PE**PF**V**DCK<u>E</u>CGRKMHQICV--<u>L</u>HYDI<u>I</u>W**P**SG--<u>FV</u>CDNC SLPFRQPVDPQLLGIPDYFDIVKNPMDLSTIKRKLDTGQYQEPWQYVDDVWLMFNNAWLYN SWPFLKPVNKKQ--VKDYYTVIKRPMDLETIGKNIEAHRYHSRAEYLADIELIATNCEQYN AWPFMEPVKRTE--APGYYEVIRSPMDLKTMSERLKNRYYVSKKLFMADLQRVFINCKEYN SWPFLKLVSKIQ--VPDYYDIIKKPIALNIIREKVNKCEYKLASEFIDDIELMFSNCFEYN IYPFHTPVNAKV--VKDYYKIITRPMDLQTLRENVRKRLYPSREEFREHLELIVKNSATYN -KSMDGDE-CSFCLG-CSSCRD-CMFCG-COIC-CKIC-U13646 MOZ.2 TFIID MOZ.1 RBBP2 PCAF p300 CCG1 CBP BAZ



Position	Distance (cRays)
D14S835 — D14S725 —	8.8
	18.9
D14S730 —— BAZ —— D14S75 ——	5.6 8.9
D14S728	12.1
D14S872	10.1

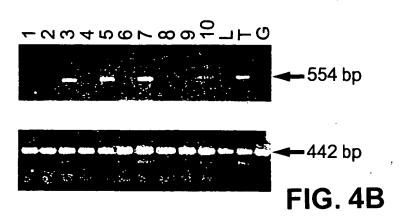
Chr 14

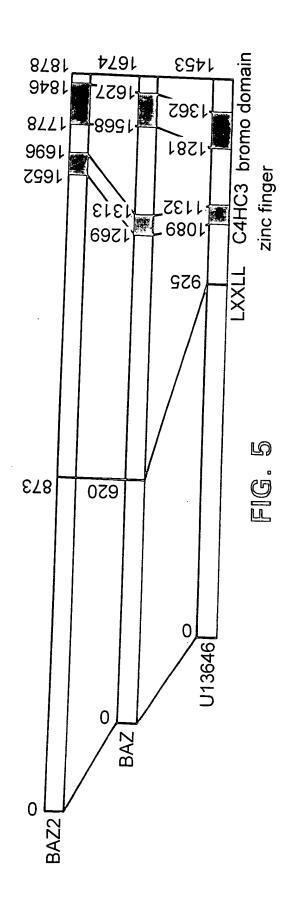
FIG. 2B



_ 2.4

FIG. 4A

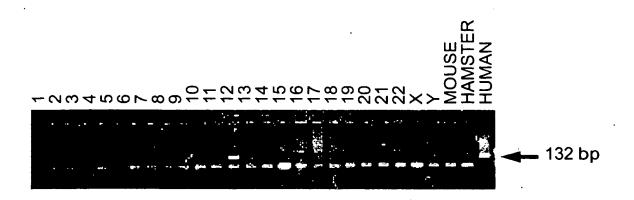


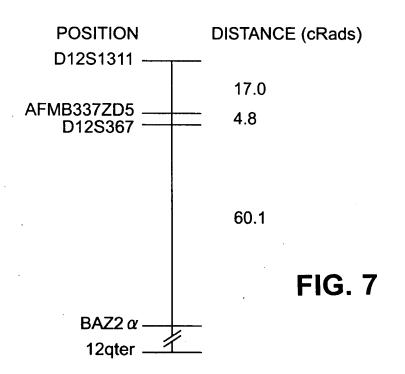


:IG. 6E

BAZ	SWPFLKLVSKIQVPDYYDIIKKPIALNIIREKVNKCEYKLASEFIDDIELMFSNCFEYN
922	SWPFHHPVNKKFVPDYYKVIVNPMDLETIRKNISKHKYQSRESFLDDVNLILANSVKYN
BAZ2	AWPFLEPVNPRLVSGYRRIIKNPMDFSTMRERLLRGGYTSSEEFAADALLVFDNCQTFN
PCAF	AWPEMEDVKRTEAPGYYEVIRSPMDLKTMSERLKNRYYVSKKLFMADLQRVFINCKEYN
U13646	U13646 ALPFLEPVNPKLVPGYKMIISKPMDLKTIROKNEKLIVSETYQFCFFAIFDLKLKMKITQYETPEDFAEDIELMFANCRQFN
CBP	SLPFRQPVDPOLLGIPDYFDIVKNPMDLSTIKRKLDTGQYQEPWQ <u>YV</u> DDVWLMFNNAWLYN
	FIG. 6A
C4H(C4HC3 Zn finger
	* * * * *
BAZ	CKICRPKIKTVPEGD-WFCPEC
013646	
RBBP2	
MOZ.1	!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
MO 7, 2	CSSCRDDPPLTRMPKGM-WICQIC
D300	GESVSLGDDPSQPQT
BAZ2	CLVCREMEAVEGD-WFCTVC

bromodomains





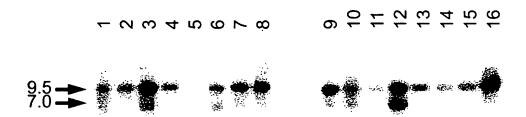
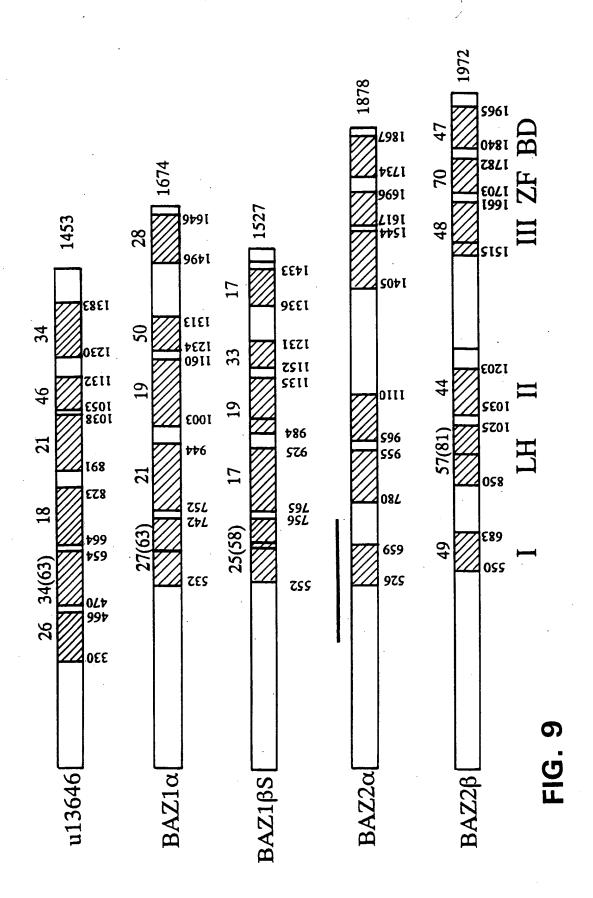


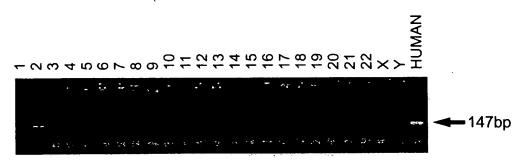
FIG. 8A

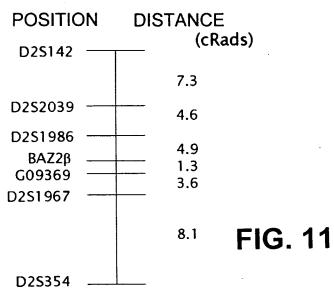


FIG. 8B



TR.LPPEIFG NQCLSSQGFA GLTLPSGAFS	GLVLSGSTFS _	LFFFLTAIFO TROLLRLALE	LVRLLKAALH LVRLLSAAVC	SANAKYRYQK	•	•		i	FIG. 10
	1 LEQRRIELEMAKELKKPNEDM.CLADOKPLPELPRIPGIVLSGSTFS	DALMVLEFL DALMVHEFV	47 DCLTIVEFLHSFGKVLGFDPAKDVPSLGVLQEGLLCQGDSLGEVQDLLVRLLKAALH 47 DCLMVVQFLRNFGKVLGFDVNIDVPNLSVLQEGLLNIGDSMGEVQDLLVRLLSAAVC	103 AIAEEEEEVAKEQLTDADTKGCSLKSLDLDSCTLSEILRLHILASGADVTSANAKYRYQK 116 FPGMGNEKRFGQGGGEMGLDRENFSEVMRLFLIDKGKR	DPGFPSYCQSLKIL	104 DPGLITGYKAKTALGEHLLNVGVNRDNVSEILQIFMEAHCGQ	163 RGGFDATDDACMELRLSNPSLVKKLSSTSVYDLTPGEKMKILHALCGKL 154GEELSQPLLTCNFLSISPEQKASILAFLCDEL	146PALCDRLRTQPFQAQPPQQKAAVLAFPVHEL	146TELTESLKTKAFQAHTPAQKASVLAFLINEL
baz1α u13646 baz2α	baz2β	baz1α u13646	baz2α baz2β	bazl α 1 u13646 1		baz2β 1	$\begin{array}{ccc} \text{bazl}\alpha & 1\\ \text{ul3646} & 1 \end{array}$	z2a	$baz2\beta$ 1
ğağ	Ą	מָׁב	ba	מ ֹח	ď	Ą	ď 'n	ba	Ą





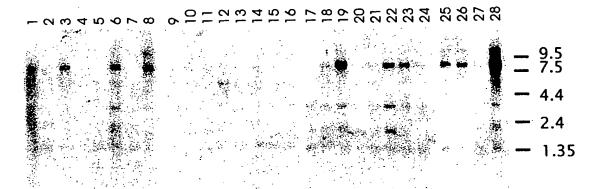


FIG. 12

> YPITAVSLMEALSADKGGFLYLNRVLVILLQTLLQTLLQDEIAEDYGELGMKLSEIPLTLHSVSE 689 -LQTLLQDEIAEDYGELGMKLSEIPLTLHSVSE 069 619 089 670 629 099 YPITAVSLMEALSADKGGFLYLNRVLVIL-650 649 BAZIβS $BAZ1\beta L$

BAZ1Bs BAZ1A BAZ2A BAZ2B	1 1	
BAZ1Bs BAZ1A BAZ2A BAZ2B	1 18	PPAPAASGLKPSPSSGEGLYTNGSPMNFPQQGKSLNGDVNVNGLSMGQTKSTSSGGGNRKCNQEQSKNQPLDARVDKIKDKKPRK.KAMESSSNSDSDS
BAZ1Bs BAZ1A BAZ2A BAZ2B	1 63	
BAZ1Bs BAZ1A BAZ2A BAZ2B	1 122	GQYPLNGILGGSRQPSSPSHNTNLRAGSQKFWANGTHSPMGLNFDSQELYDSFPDQNFE. NQVLLHGISDPKADGQKATEKAQEKRIHQPLPLAFESQTHSFQSQQKQP
BAZ1Bs BAZ1A BAZ2A BAZ2B	1 181	MEDASESSRGVAPLINNVVLPGSPLSLPVSVTGCKSHRVANKKVEARSEKLLPTALPPSE EVCSGIHPDEAAEKEMTSVVAENGTGLVCSLELEEEQPELKMCGYNGS QVLSQQLPFIFQSSQAKEESVNKHTSVIQSTGLVSNVKPLSLVNQAKKETYM
BAZ1Bs BAZ1A BAZ2A BAZ2B	61 229	PKVDQKLPRSSERRGSGGGTQFPARSRAVAAGEAAARGAAGPERGSPLGRRVSPRCLCSG VPSVESLHQEVSVLVPDPTVSCLDDPSHLPDQLEDTPILSEDSLEPFNSLAPEPVSG KLIVPSPDVLKAGNKNTSEESSLLTSELRSKREQYKQAFP
BAZ1Bs BAZ1A BAZ2A BAZ2B	121	MAPLLGRKPFPLVNPLPGEEPFFTIPHT EGGQVAVGVIAGKRGRRGRDGSRRAPGGREMPLLHRKPFVRQKPPADLRPDEEVFYCKVT GLYGIDDTELMGAEDKLPLEDSPVISALDCPSLNNATAFSLLADDSQTSTSIFASP.TSPSQLKKQESSKSLKKVIAALSNPKATSSSPAHPKQTLENNHPNPFLTN
BAZ1Bs BAZ1A BAZ2A BAZ2B	181 345	QEAFRTREEYEARLERYSERIWTCKSTGSSQLTHKEAWEEEQEVAELLKEEFPAWYEKLV NEIFRHYDDFFERTILCNSLVWSCAVTGRPGLTYQEALESEKKARQNLQS.FPEPLI PVLGESVLQDNSFDLNNGSDAEQEEMETQSSDFPPSLTQPAPDQSSTIQLHP ALLGNHQPNGVIQSVIQEAPLALTTKTKMQSKINENIAAASSTPFSSPV.NLSTSGRRTP
BAZ1Bs BAZ1A BAZ2A BAZ2B	237 397	LEMVHHNTASLEKLVDTAWLEIMTKYAVGEECDFEVGKEKMLKVKIVKIHPLEKVD IPVLYLTSLTHRSRLHEICDDIFAYVKDRYFVEETVEVIRNNGARLQCTILEVLP ATSPAVSPTTSPAVSLVVSPAASPEISPEVCPAASTVVSPAVFSVV GNQTPVMPSASPILHSQGKEKAVSNNVNPVKTQHHSHPAKSLVEQFRGTDSDIPSSK

BAZ1Bs BAZ1A	292	EEATEKKSDGACDSPSSDKENSSQIAQDHQKKETVVKEDEGRRESINDRARRSPRKLPSHQNGFANGHVNSVDGETIIISDSDDSETQS
BAZ2A BAZ2B	443 407	<u>D</u> SEDSNED <u>E</u> EEDDEEEDEEDDEDDESDDSQSESDSNSESD <u>TE</u> GSEE E DDDD <u>K</u> DQDESDSD
BAZ1Bs BAZ1A	202 324	PTSLKKGERKWAPPKFLPHKYDVK.LQNEDKIISNVPADSLIRTERPPNKEIVRYFTRHN.CSFQNGKKKDAIDPLL.FKYKVQPTKKELHESAIVKATQISRRKHLFSRDKLKLFLKQH
BAZIA BAZZA	443	SPASSAVLPAVSLEVPLTASVTSPKASPVTSPAAAFP.TASPANKDVSS
BAZ2B	467	TEGEKTSMKLNKTTSSVKSPSMSLTGHSTPRNLHIAKAPGSAPAALCSESQSPA
BAZ1Bs		ALRAGTGENAPWVVEDELVKKYSLPSKFSDFLLDPYKYMTLNPSTKRKNTGSPD
BAZ1A BAZ2A		CEPQEGVIKIKASSLSTYKIAEQDFSYFFPDDPPTFIFSPANRRRGRPP FLETTADVEEITGEGLTASG.SGDVMRRRIATPEEVRLPLQHGWRREVRIKKGSHRWQGE
BAZZB		FLGTSSSTLTSSPHSGTSKRRRVTDERELRIPLEYGWQRETRIRNFGGRLQGE
BAZ1Bs	315	RKPSKKSKT DN SSLSSPLNPKLWCH VH LKKSLSGSPLKVKNSKNSKSPEEHLEEMMKMMS
BAZ1A	431	KRI.HISQEDNVANKQTLASYRSKATKER
BAZ2A BAZ2B	550	TWYYGPCGKRMKQFPEVIKYLSRNLVHSVRREHFSFSPRMPVGDFFEERDTPEGLQWVQL
BAZZB	5/4	VAYYAPCGKKLRQYPEVIKYLSRNGIMDISRDNFSFSAKIRVGDFYEARDGPQEMQWCLL
BAZ1Bs	375	PNKLHTNFHIPKK G PP <u>AK</u> K P GKHS DK P L KAKGRSK G ILNGQ K STGNSKSP KK G LK TP K TK
BAZ1A	459	QEEMKSLAFEKAK
BAZ2A	610	SAEEIPSRIQAITGKRGRPRNTEKAKTKEVPKVKRGRGRPPKVKITELLNKTDNRPLKKL
BAZ2B	634	KEE <u>DV</u> IPRIRAMEGRRGRPPNP <u>DRQR</u> AREES <u>RMR</u> RR <u>K</u> GRPPNVGNAEFL <u>D</u> NADAKLL <u>R</u> KL
BAZ1Bs	435	MKQMTLLDMAKGTQKMTRAPRNSGGTPRTSSKPHKHLPPAALHLIAYYKENKDREDKRSA
BAZ1A	477	LKREKADALEAKKKEKEDKEKKR
BAZ2A	670	EAQETLNEEDKAKIAKSKKKMRQKVQRGECLTTIQGQARNKR
BAZ2B	694	QA QE IARQA A QIKLLRK L QKQEQ A RV AK EA KK QQAI <u>M</u> AAEE K RKQK E QIKI <u>M</u> KQQE K I K R
BAZ1Bs	495	LSCVISKTARLLSSEDRARLPEELRSLVQKRYELLEHKKRWASM
BAZ1A	500	<u>E</u> ELKK <u>I</u> VEEER <u>L</u>
BAZ2A	712	KQETKSLKHKVKREKKEKUK
BAZ2B	754	<u>IQQIR</u> MEKELRAQQIL EAKKK KKE E AAN <u>A</u> KLL E AEK <u>RI</u> K E KEMRRQQAVL <u>L</u> K <u>H</u> QE <u>R</u> E <u>R</u> R <u>R</u>
		L L L L
		SEEQRKEYLKKKREELKKKLKEKAKERREKEMLERLEKQKR.YEDQEL
		KKKEEKERLKVEREKEREKLREEKRKYVEYLKQWSKPREDMEC
		MKEKEEVTKAKPACKADK.TLATORRLEEROKOOMILEEMKKPTEDMCL
BAZ2B	014	QHMMLMKAMEARKKAEEKERLKQEKRDEK.RLNKERKLEQRRLELEMAKELKKPNEDMCL
		L L L L LL LL LL
		TG.KNLPAFRLVDTPEGLPNTLFGDVAMVVEFLSCYSGLLLPDAQYPITAVSLMEAL.
BAZ1A		DDLKELPEPTPVKTRLPPEIFGDALMVLEFLNAFGELFDLQDEFPDGVTLEVLEEALV
BAZ2A BAZ2B		TDHQPLPDFSRVPGLT.LPSGAFSDCLTIVEFLHSFGKVLGFDPAK.DVPSLGVLQEGLL ADQKPLPELPRIPGLV.LSGSTFSDCLMVVQFLRNFGKVLGFDVNI.DVPNLSVLQEGLL
DMUZD	ψIJ	UDSVEHEBRI VIEGRA - FOGO LE ODCIMIA A ĀLITVINE GVA FIGERA MIT - DA KIATOA FIÑE OPP

	LXXLL L LXXLLXXLL
BAZ1Bs 64	2 S.ADKGGFLYLNRVLVILLQTLLQDEIAEDYGELGMKLSEIPLTLHSVSE
	G.NDSEGPLCELLFFFLTAIFQAIAEEEEEVAKEQLTDADTKGCSLKSLDLDSCTLSE
BAZZA 86	
BAZ2B 93	1 NIGDSMGEVQDLLVRLLSAAVCDPGLITGYKAKTALGEHLLNVGVNRDNVSE
	L L L
BAZ1Bs 69	
BAZ1A 67	
BAZZA 91	
BAZ2B 98	3 ILQIFMEAHCGQTELTESLKTKAFQAHTPA
	L L LL
BN71Bc 74.	E KLQILTALCHRILMTYSVQD
	D EKMKILHALCGKLLTLVSTRDFIEDYVDILRQAKQEFRELKAEQHRKEREEAAARIRKRK
	3 QKAAVLAFPVHELNGSTLIINEIDKTLESMSSYRKNKWIVEGRLRR
	3 QKASVLAFLINELACSKSVVSEIDKNIDYMSNLRRDKWVVEGKLRK.
D.1022 102	2-110-1-11-1-1-1-1-1-1-1-1-1-1-1-1-1-1-1
BAZ1Bs 78	LAV LKE END KK RAEKQ K R KEME AK N K E NGKVENGL G KTD R K K RIVKF E PQV DTE AE
BAZ1A 79	D EEKLKEQEQKMKEKQEKLKEDEQRNSTADISIGEEEREDFDTSIESK.DTEQKELDQ
BAZ2A 98	
BAZ2B 105	OLRIIHAKKTGKRDTSGGIDLGEEQHPLGTPTPGRKRRKGGDS.DYDDDDDDDDDDDD
	DMISAVKSRRLLAIQAK.KEREIQEREMKVKLERQAEEERIRKHKAAAEK
	5 DMFTEDEDDPGSHKRGRRGKRGQNGFKEFTRQEQINCVTRELLTADEEEALKQEHQRKEK
	GMEEEEEESIAAVPGRRGRRDGEVDATASSIPELERQIEKLSKRQLF
BAZZB III	GDEDDEDEEDKEDQKGKKTDICEDEDEGDQAASVEELEKQIEKLSKQQSQ
BAZ1Bs 88	AFQEGIAKAKLVMRRTPIGTDRNHNRYWLFSDEVPGLFIEKGWV.HDSIDYRFNHHC
	ELLEKIQSAIACTNIFPLGRDRMYRRYWIFPS.IPGLFIEEDYSGLT.EDMLLPRPSSF.
BAZ2A 107	
BAZ2B 116	
	LXXLL
	KDHTVSGDEDYCPRSKKANLGKNASMNTQHGTATE
	3 QNNVQSQDPQVSTKTGEPLMSESTS.NIDQGPRDH
	VAAHASLNPALFSMKMELAGSNTTASSPARARSRP.LKTKPGFMQ.PRHFKSPVRGQ
BAZ2B 122	KAESVQIKEEMFETSGDSLNCSNTDHCEQKEDLKEKDNTNLFLQKPGSFSKLSKLL
BA71Be 97	7
	7
	DSEQPQAQLQPEAQLHVPAQPQPQLQLQLQSHKGFLEQEGSPLSLGQSQHDLSQSAFLSW
	B EV AKMPPESEVMTP. KPNAGANGCTLSYQNSGKHSLGSVQSTATQSN
	- - - - -
	1
	7
	LSQTQSHSSLLSSSVLTPDSSPGKL.DPAPSQPPEEPEPDEAESSPDLQAFWFNISAQMP
BAZ2B 132	VEKADSNNLFNTGSSGPGKFYSPLPNDQLLKTLTEKNRQWFSLLPRTP

	CNAAPTPPLAVSEDQPTPSPQQLASSKPMNRPSAANPCSPVQFSS. CDDTSLTHADMSTASLVTPQSQPPSKSPSPTPAPLGSSAQNPVGLNPFALSPLQVKGG
BAZ1A 997 BAZ2A 1350	TPLAGLAPKRRAGDPGEMPQSPTGLGQPKRRGRPPSKFFKQMEQRYLTQ VSMMGLQFCGWPTGVVTSNIPFTLSVPSLGSGLGLSEGNGNSFLTS
BAZ1A 997 BAZ2A 1399	LXXLLVAVETTTPKQGQNLWFLCDSQKELDELLSVQLPKPVHKPNRWCFYSSCEQLDQLI LTAQPVPPEMCSGWWMIPDPEMLDAML NVASSKSESPVPQNEKATSAQPAAVEVAKPVDFPSPKPIPEEMQFGWWRIIDPEDLKALL
BAZ1A 1024 BAZ2A 1426	NCLHPQGIRESQLKERLEKRYQDIIHSIHLARKPNLGLKSEALNSRGHRESALKETLLQEKSRICAQLARFSEEKFHFSDKPQPDSKPTYSRGRSSNAYDKALHPRGIREKALHKHLNKHRDFLQEVCLRPSADPIFKVLHLRGIREKALQKQIQKHLDYITQACLKNKDVAII
BAZ1A 1084 BAZ2A 1463	CDGNQELLNFLRSDLIEVATRLQKGGLGYVEETSEFEARVISLEKLKDFGECVIAL PSQMCAEKQLELR.LRDFLLDIEDRIYQGTLGAIKVTDRHIWR EPRQLPAFQEGIMSWSPKEKTYETDLAVLQWVEELEQRVIMSDLQIRGWTCPSPD ELNENEENQVTRDIVENWSVEEQAMEMDLSVLQQVEDLERRVASASLQVKGWMCPEPA
BAZ1A 1126 BAZ2A 1518	QASVIKKFLQGFMAPKQKRRKLQSEDSAKTEEVDEEKKMVEEAKV
BAZ1A 1183 BAZ2A 1566	
BAZ1A 1243 BAZ2A 1626	C C C C H C TFSRMHVLLGMLDACIKWDMSAENARCKVCPKKGEDDKLILCDECNKAFHLFCLRPALYE SLSQVFLHLSTLDRSVIWSKSILNARCKICRKKGDAENMVLCDGCDRGHHTYCVRPKLKT SAAHVCLCLGHLERSIAWEKSVNKVTCLVCRKGDNDEFLLLCDGCDRGCHIYCHRPKMEA SAAQVALCIQQLQKSIAWEKSIMKVYCQICRKGDNEELLLLCDGCDKGCHTYCHRPKITT
BAZ1A 1303 BAZ2A 1686	C C VPDGEWQCPACQPAT.ARRNS.RGRNYTE.ESASEDSEDDESDEEEEEEEEEE VPEGDWFCPECRPKQRCRRLSFRQRPSLESDEDVEDSMGGEDDEVDGDEEEGQSEEEEYE VPEGDWFCTVC

BAZ1A 1363 BAZ2A 1697	PRKTIRGKHSVIPPAARSGRRPGKKPHSTRRSQP VEQDEDDSQEEEEVSLPKRGRPQVRLPVKT.RGKLSSSFSSRGQQQEPGRYPSRSQQSTPLAQQVEGEFTQKPGFPKRGQKIAKASGQTLKIKKLHVKGKKTNESKKGKK
BAZ1A 1422 BAZ2A 1718	K APPVDDAEVDELVLQTK
BAZ2A 1734	RKSANNTPENSPNFPNFRVIATKSSEQSRSVNIASKLSLQESESKRRCRKRQSPEPSPVT
	Bromodomain motif
BAZ1A 1539 BAZ2A 1771	RSSRRQSLELQKCEEILHKIVKYRFSWPFREPVTRDEAEDYYDVITHPMDFQTVQ LGRRSSGRQGGVHELSAFEQLVVELVRHDDSWPFLKLVSKIQVPDYYDIIKKPIALNIIRDLTFCEIILMEMESHDAAWPFLEPVNPRLVSGYRRIIKNPMDFSTMRDLALCSMILTEMETHEDAWPFLLPVNLKLVPGYKKVIKKPMDFSTIR
BAZ1A 1599 BAZ2A 1818	LXXL NKCSCGSYRSVQEFLTDMKQVFTNAEVYNCRGSHVLSCMVKTEQCLVVL EKVNKCEYKLASEFIDDIELMFSNCFEYNPRNTSEAKAGTRLQAFEHIQAQKLG ERLLRGGYTSSEEFAADALLVFDNCQTFNEDDSEVGKAGHIMRRFFESRWEEFYQGKQAN EKLSSGQYPNLETFALDVRLVFDNCETFNEDDSDIGRAGHNMRKYFEKKWTDTFKVS
BAZ1A 1653 BAZ2A 1878	LHKHLPGHPYVRRKRKKFPDRLAEDEGDSEPEAVGQSRDEDRRSREAEIQEWLQ LHVTPSNVDQVSTPPAAKKSRI L
BAZ1Bs1494 BAZ1A 1675 BAZ2A 1879 BAZ2B 1973	DTSLYSAKINSKDHNCFMMLVNTQFCMALTDTVT FIG. 18



FIG. 19A

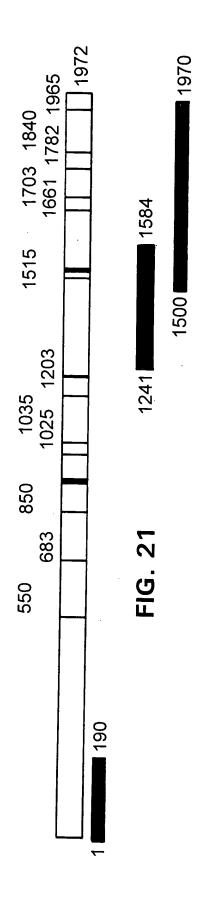
Position	Distance (cRads)
D7S1776	10.2
D7S653	8.9
D7S489 ——	7.2
BAZ1ß	16.0
D7S669	11.3
D7S675	7.6
AFM312XB5 —	<u></u>

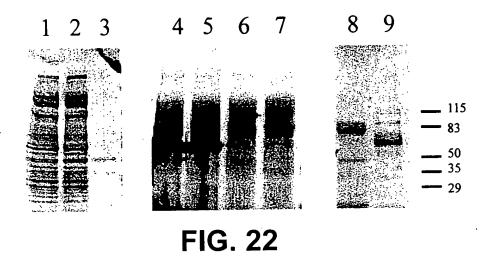
FIG. 19B

→ 7.5 kb

FIG. 20A

FIG. 20B





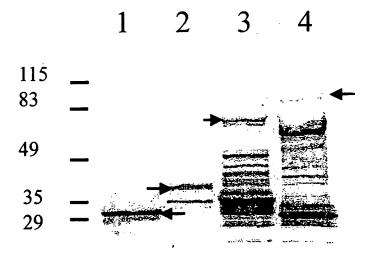


FIG. 23